

# Build Better Biologics with Machine Learning and Synbio

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April 10, 2019

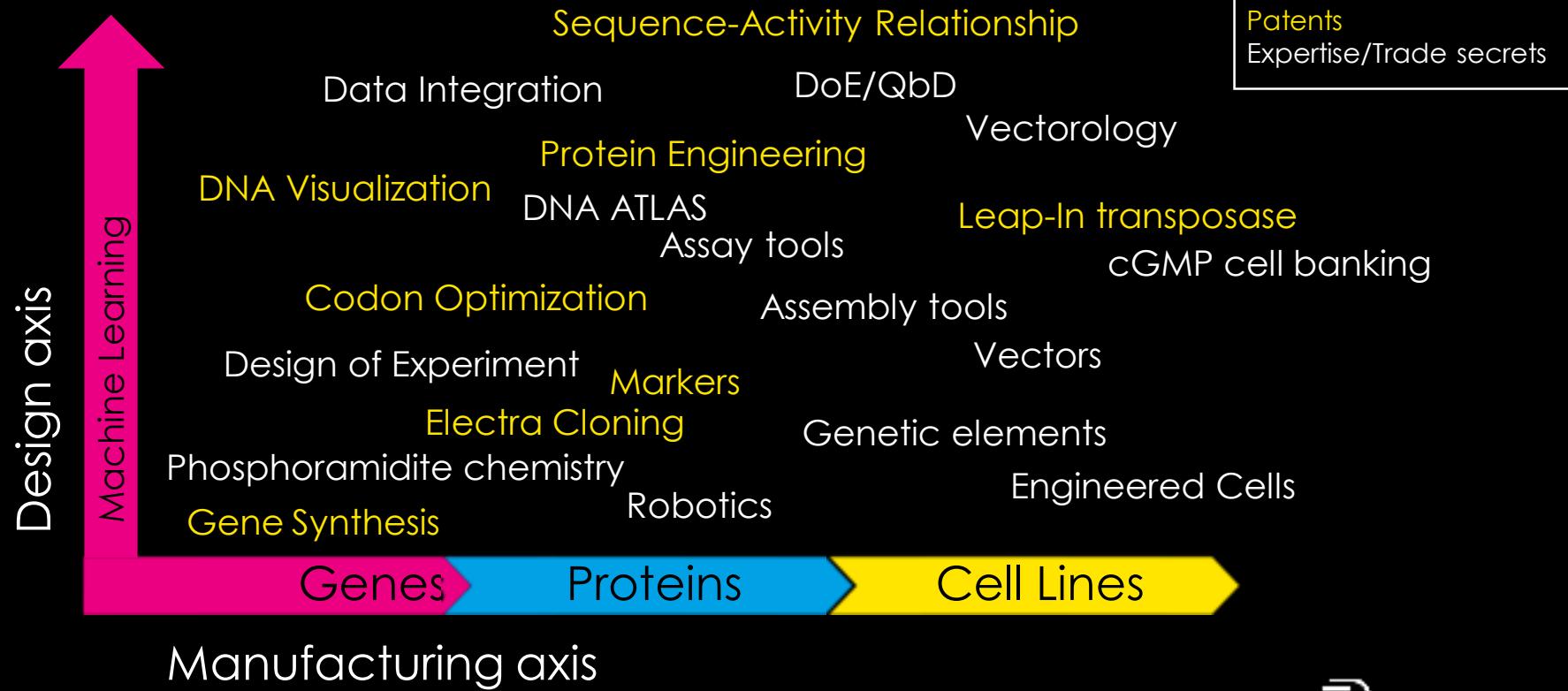




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- Founded in 2003
  - Based in SF Bay Area
  - ~100 Employees
  - 23 issued patents
  - >60 peer-reviewed papers
  - Rebranded to **ATUM** 2016



# Ecosystem of Integrated Tools



# High Throughput Biologics ATUMation

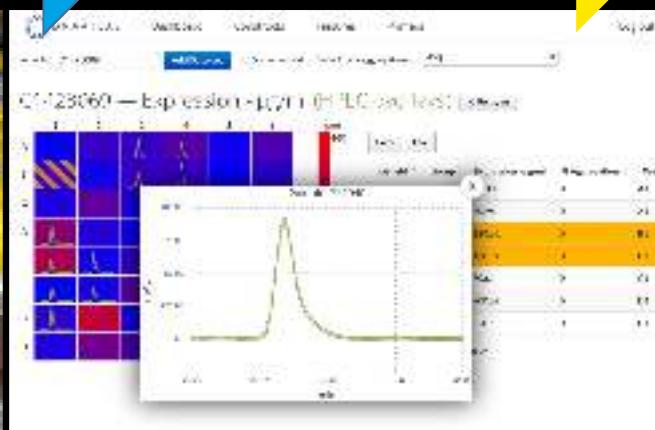
Genes



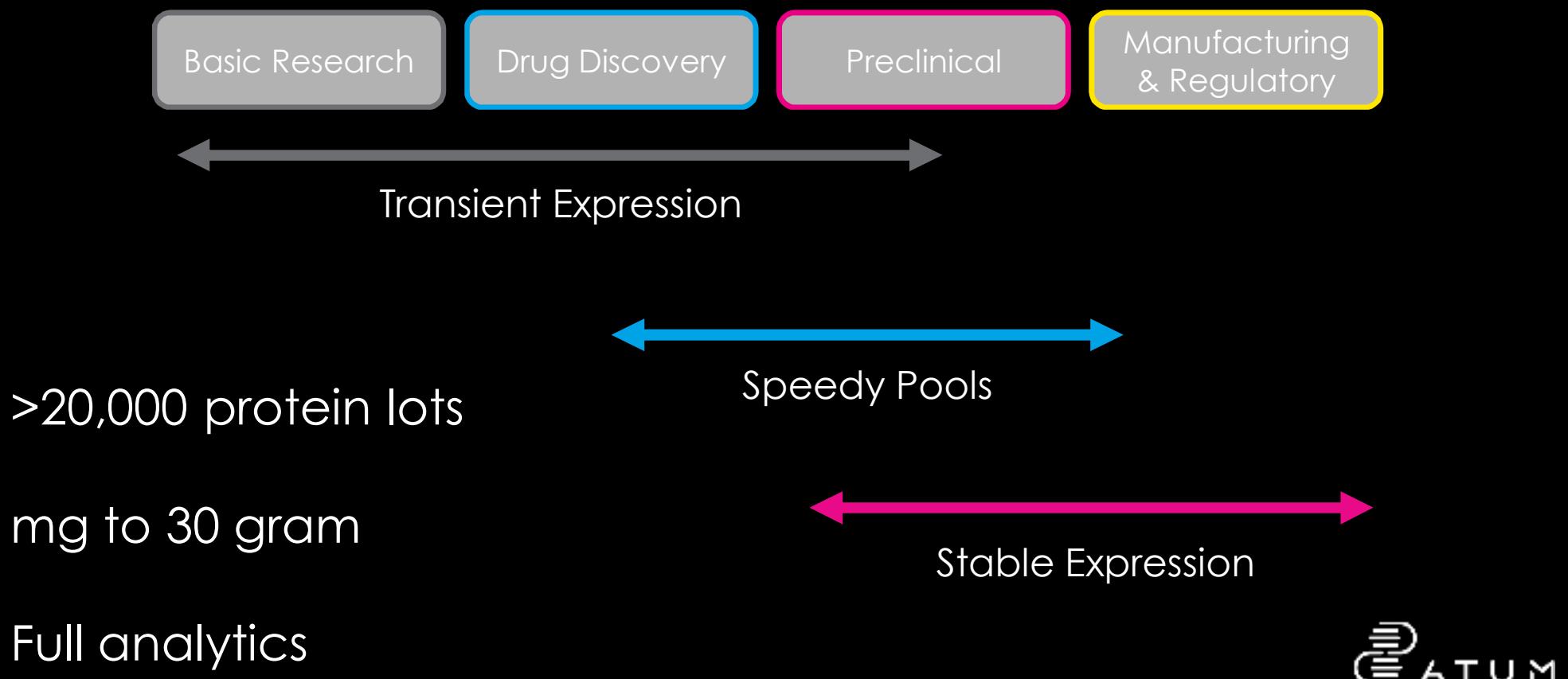
Proteins



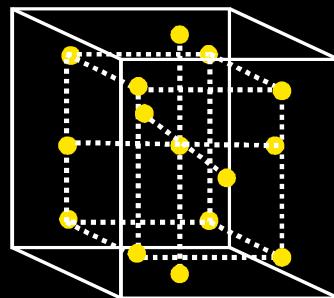
Cell lines



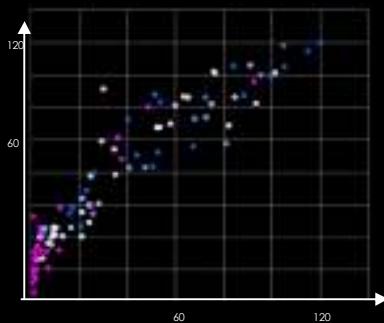
# Mammalian Protein Expression



## Design of Experiment



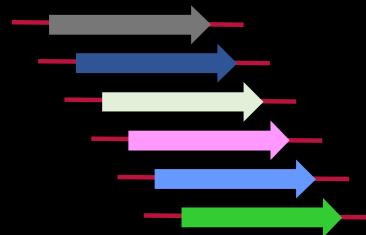
Genomic  
Data



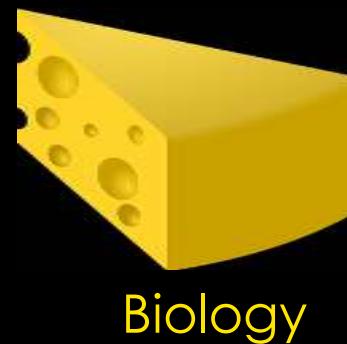
Machine  
Learning



Synthetic  
Biology

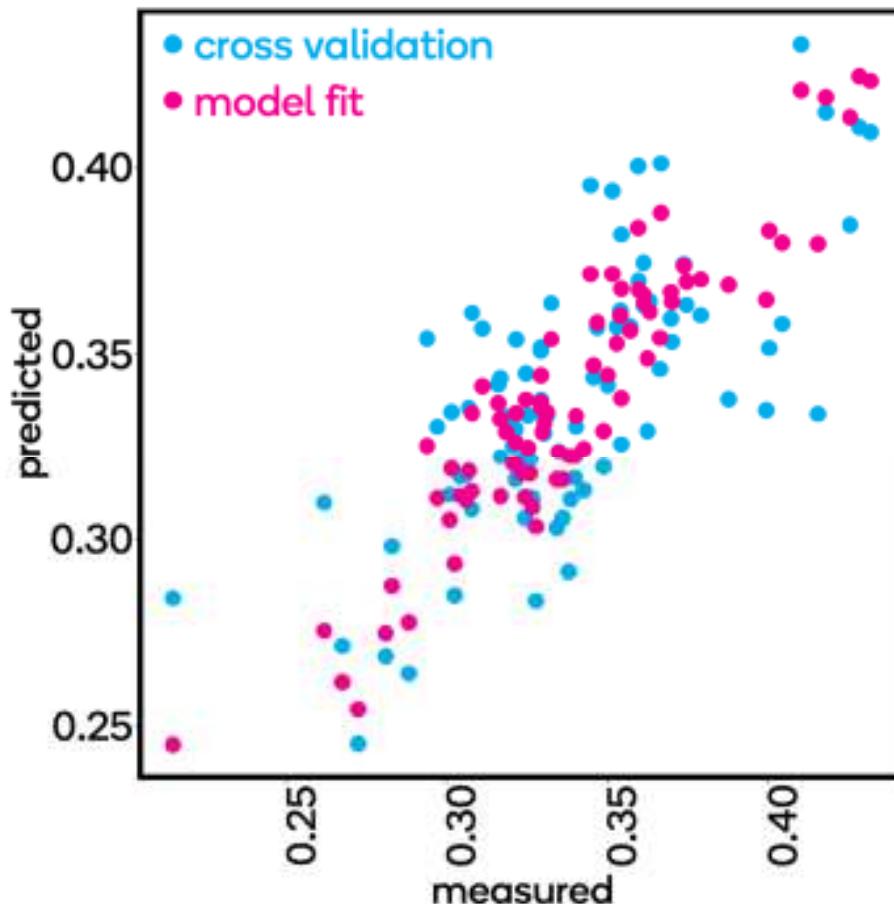


Drug  
Candidate



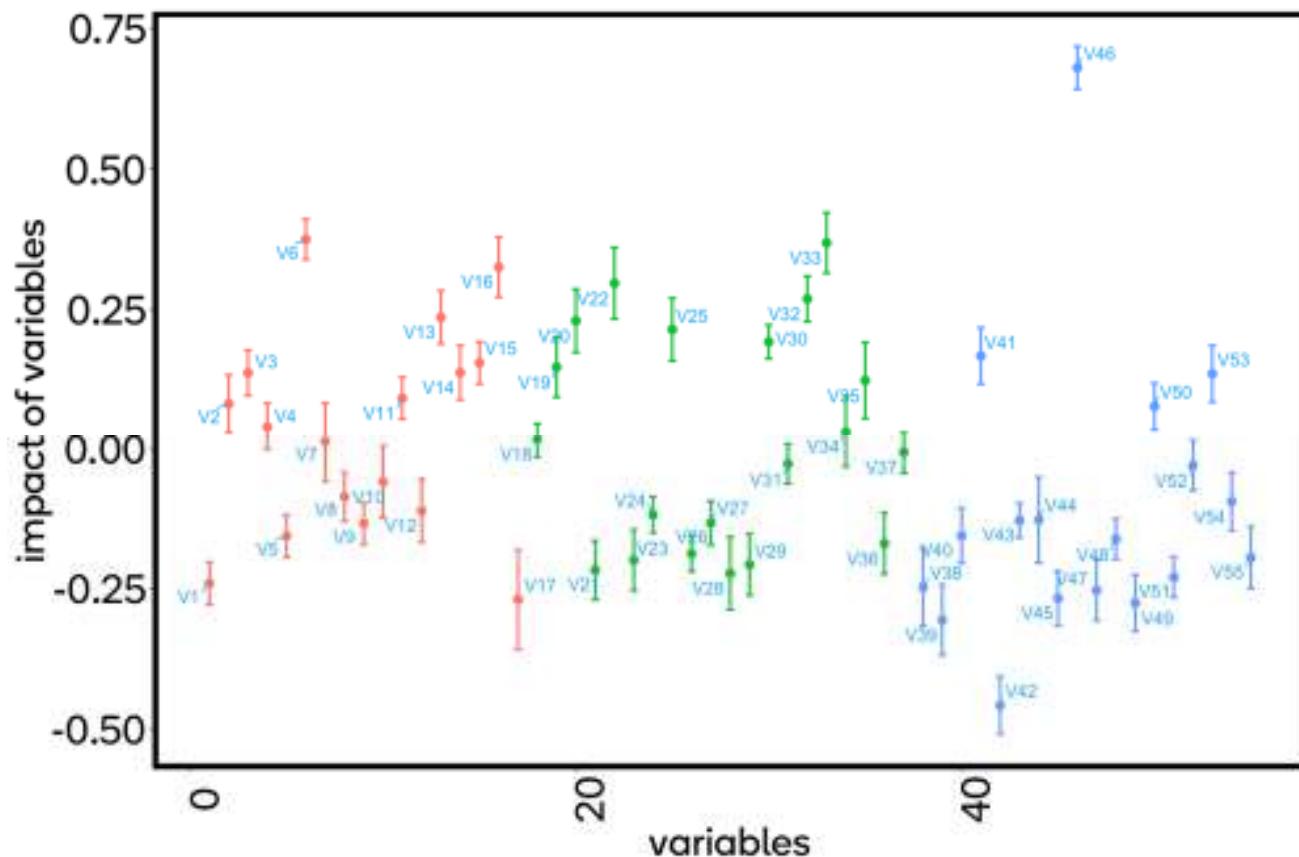
ATUM

# Machine Learning Models



- Consensus Models from several algorithms
- Models for every measured attribute
- Models on fitness functions
- Cross-validated leaving 15% out
- Only ~96 variants per round

# Machine Learning Models



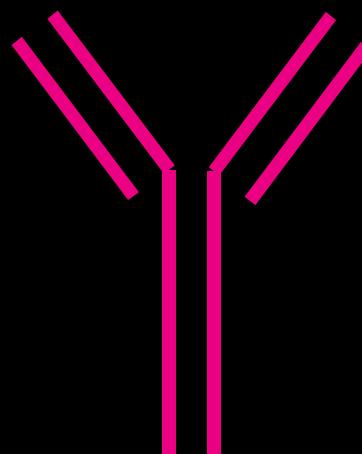
Captures additivity and epistasis  
Impact of substitutions quantified.  
~ $10^{20}$  Sequence Space explored

# Case Study: Testing 184 Variants To Capture the Space of $10^{23}$ Variants

Total available space:

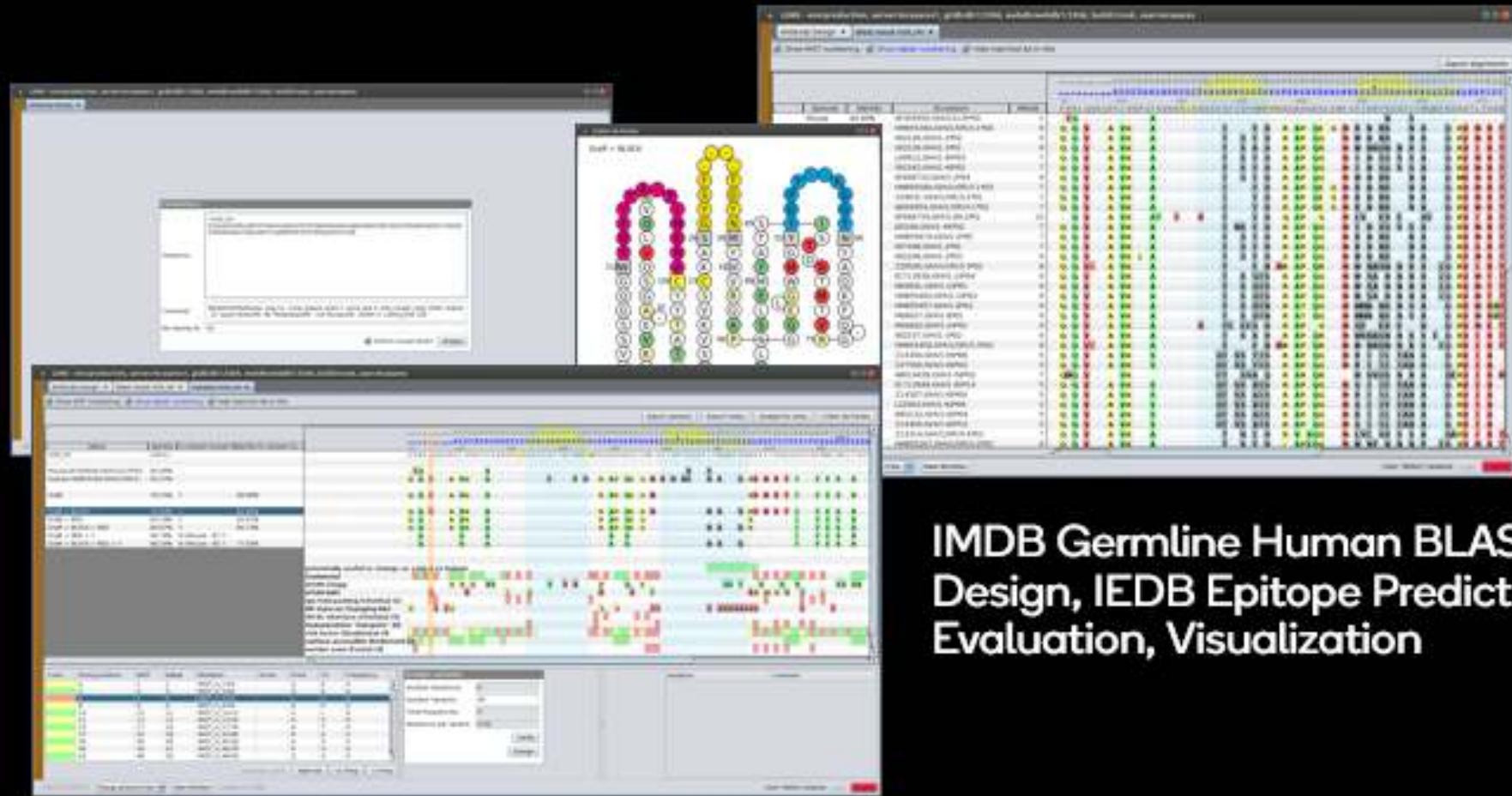
$$VH = 2^{41} (\sim 10^{12})$$

$$VL = 2^{36} (\sim 10^{10})$$



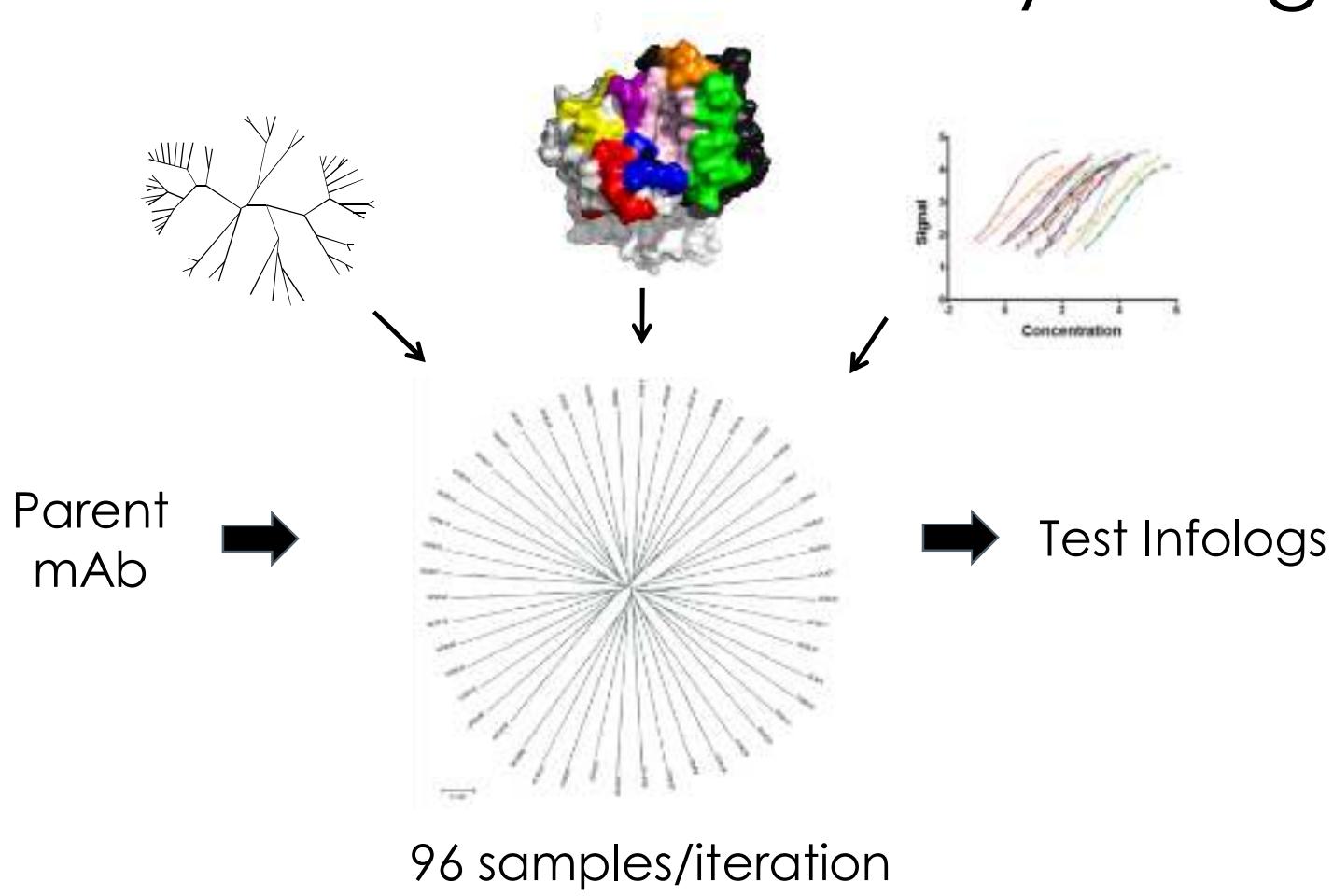
Starting from commercial molecule

# Biotech is Data Management

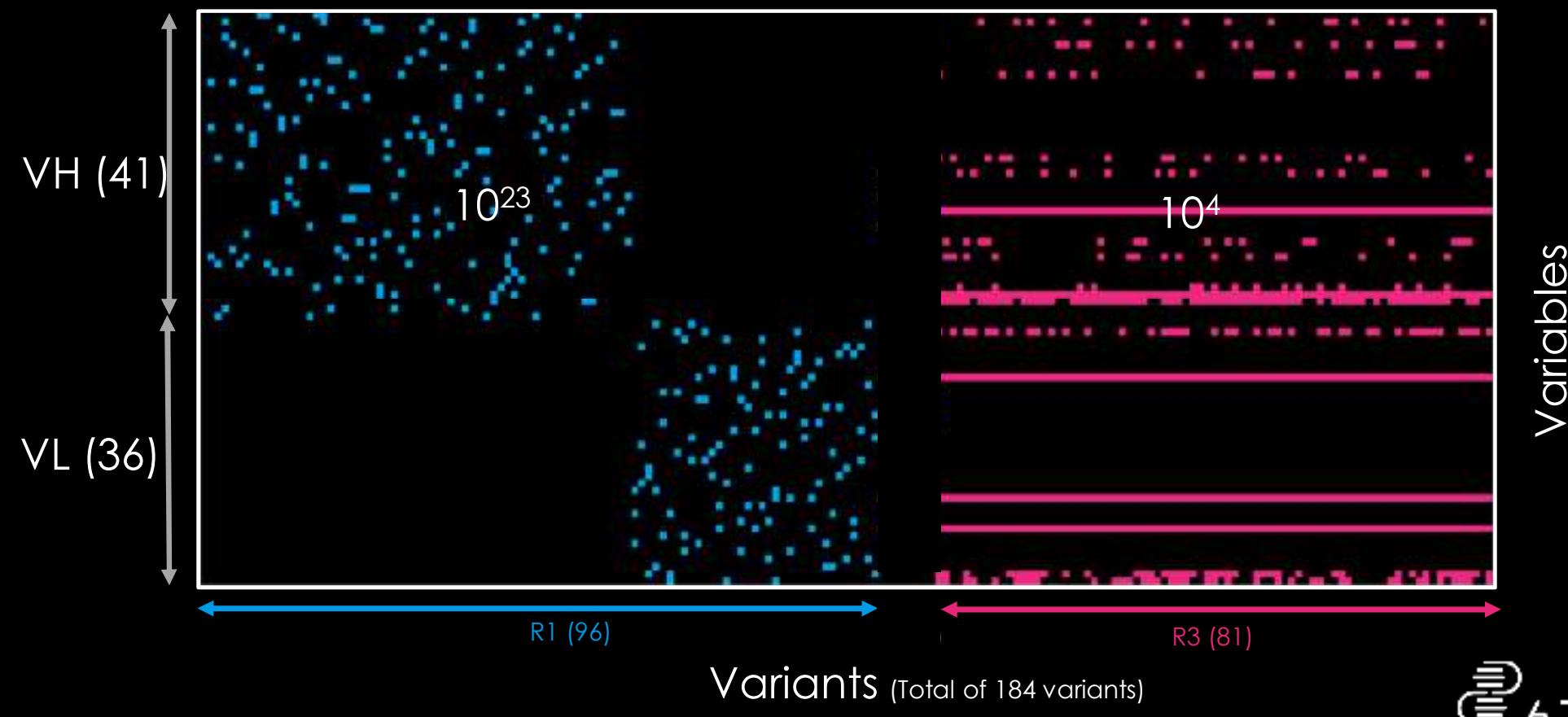


## IMDB Germline Human BLAST, DOE Design, IEDB Epitope Prediction, Evaluation, Visualization

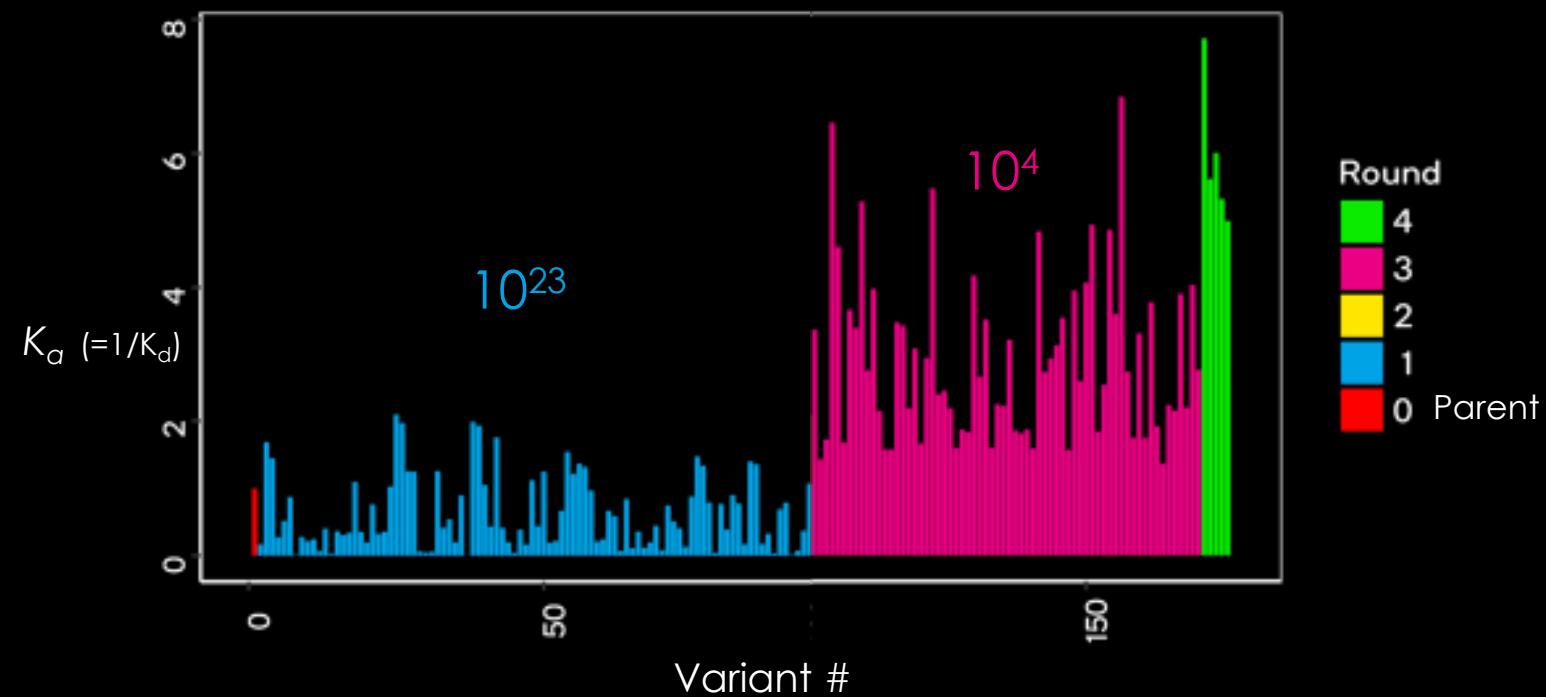
# Variable Selection and Library Design



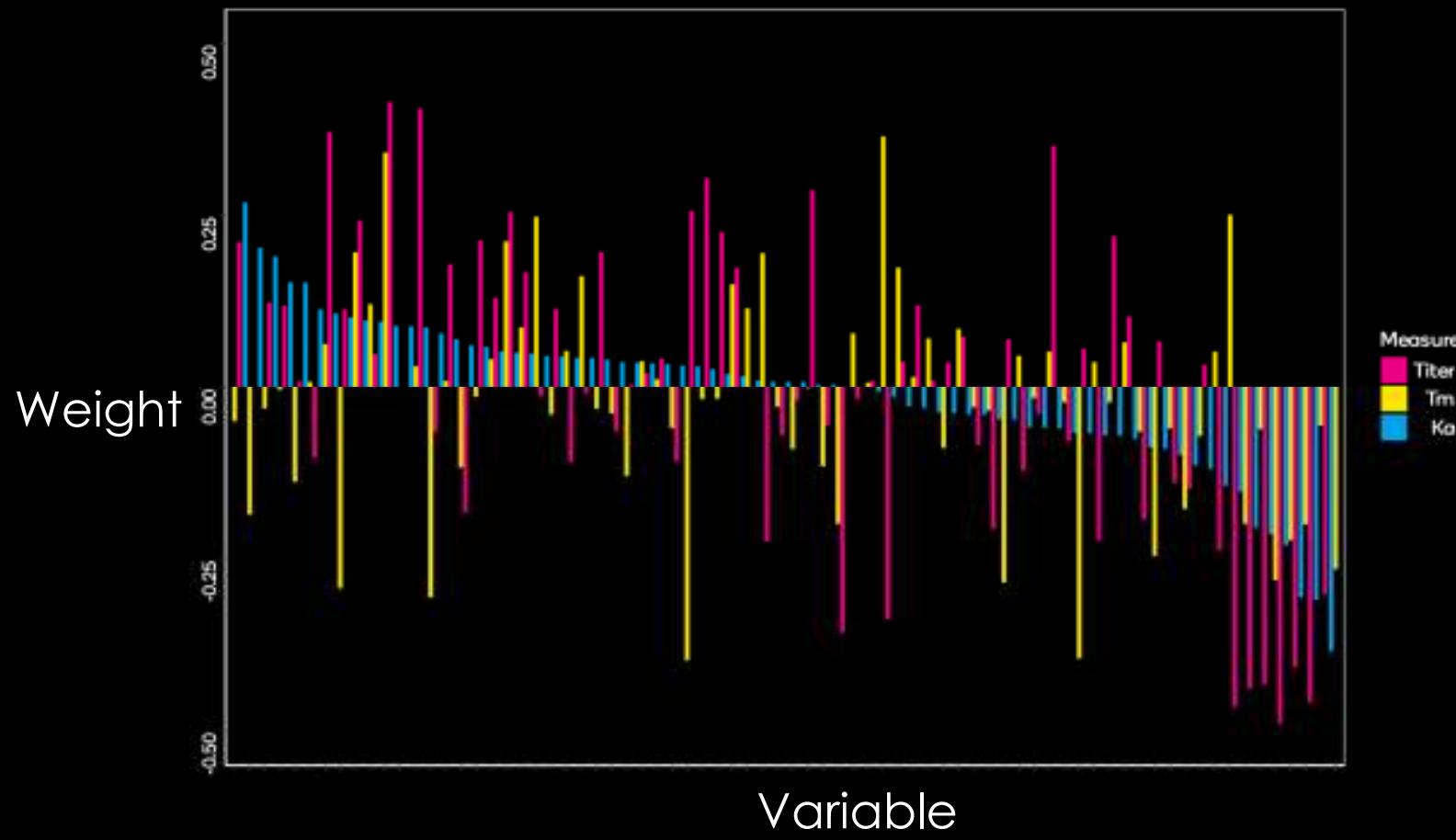
# Variable Matrix



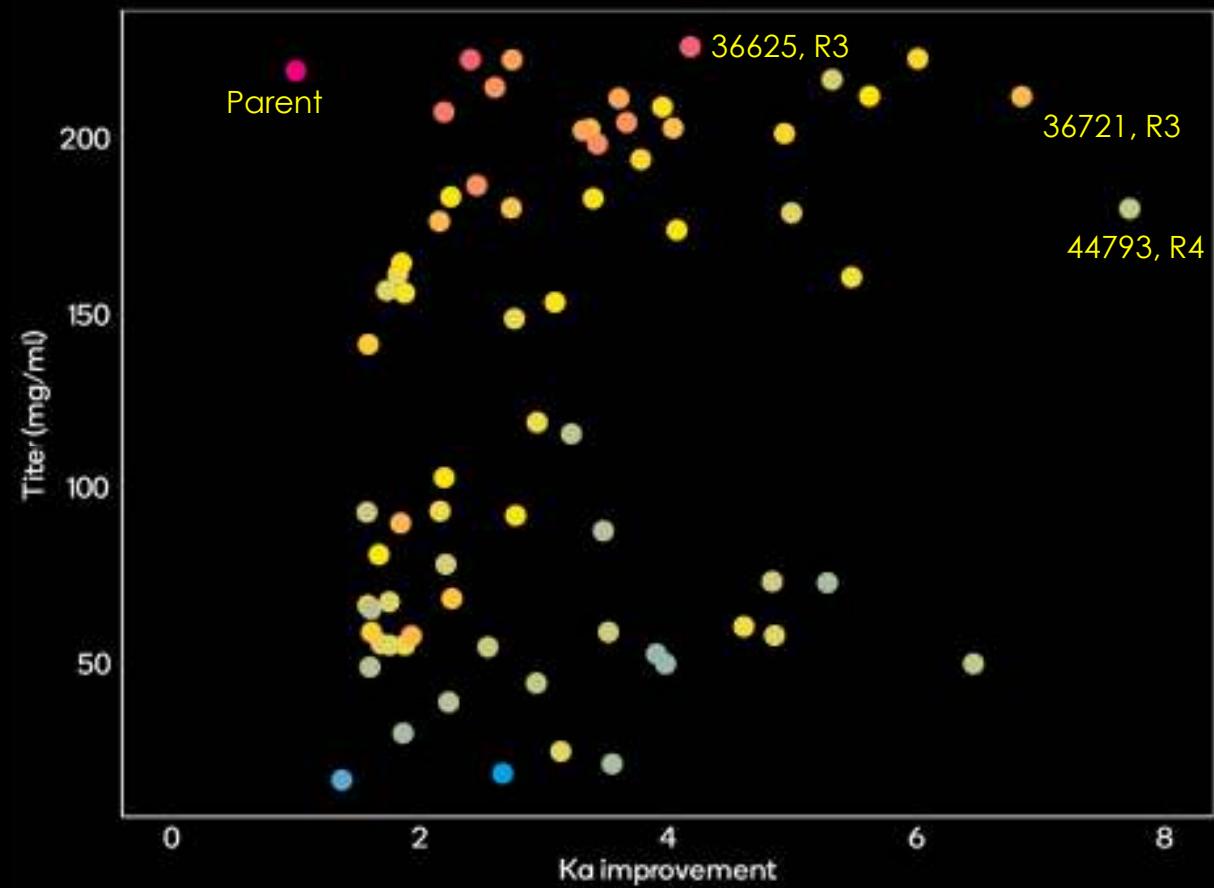
# $K_a$ Improvement by Round



# Multidimensional Variable Weight Plot



# Multi-Dimensional Improvements

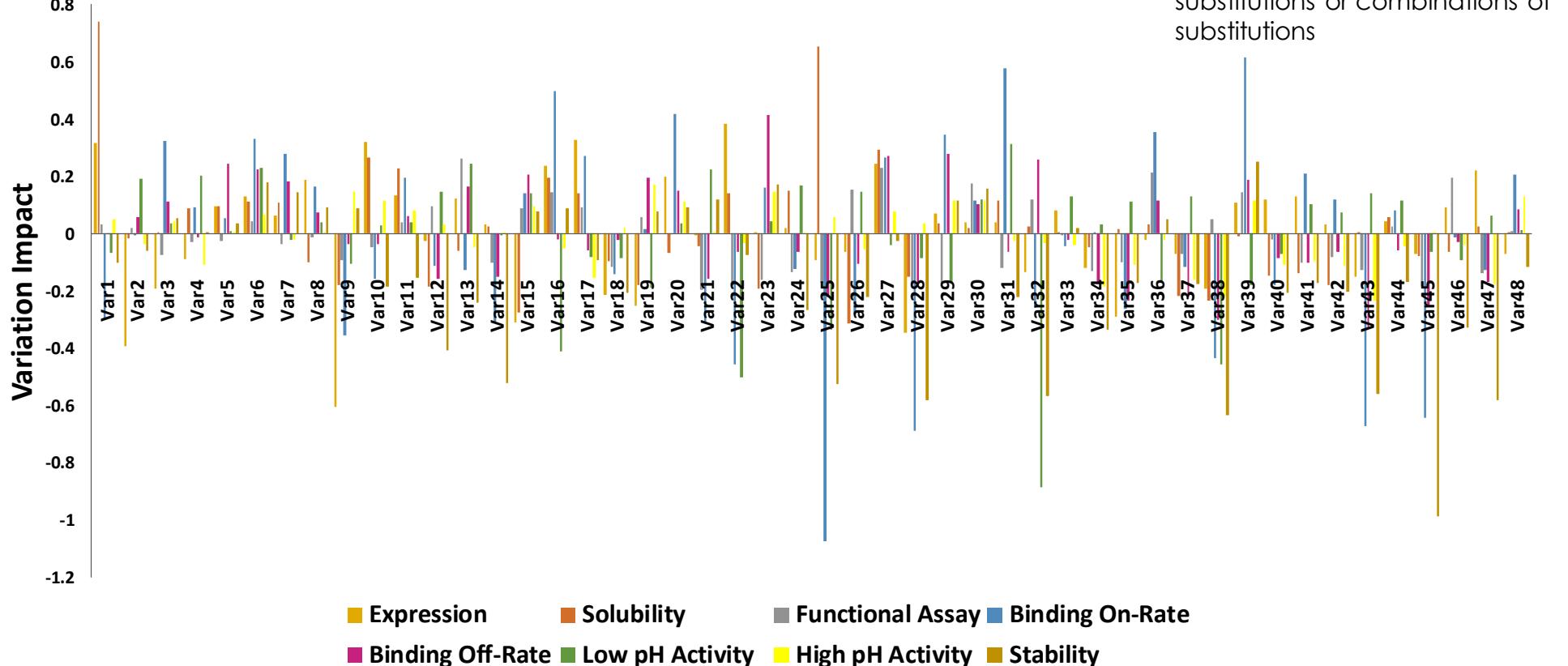


| Clone  | Titer (mg/L) | Tm (°C) | Kd (nM) |
|--------|--------------|---------|---------|
| 36721  | 212          | 59      | 0.16    |
| 36625  | 226          | 62      | 0.25    |
| 44793  | 180          | 56      | 0.14    |
| Parent | 219          | 63      | 0.98    |

Tm (°C):

- 55.0
- 57.5
- 60.0
- 62.5

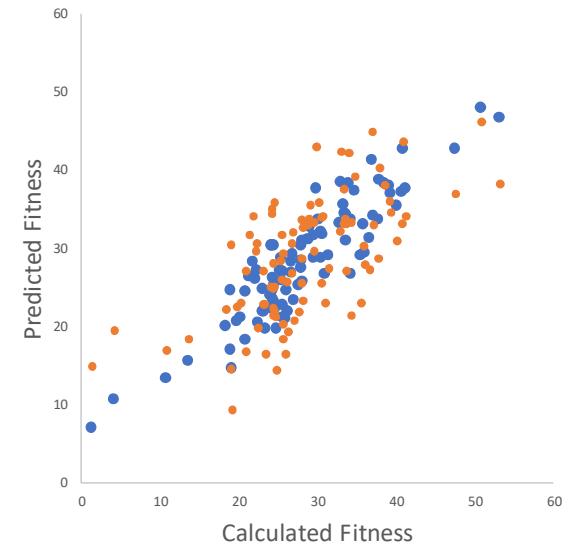
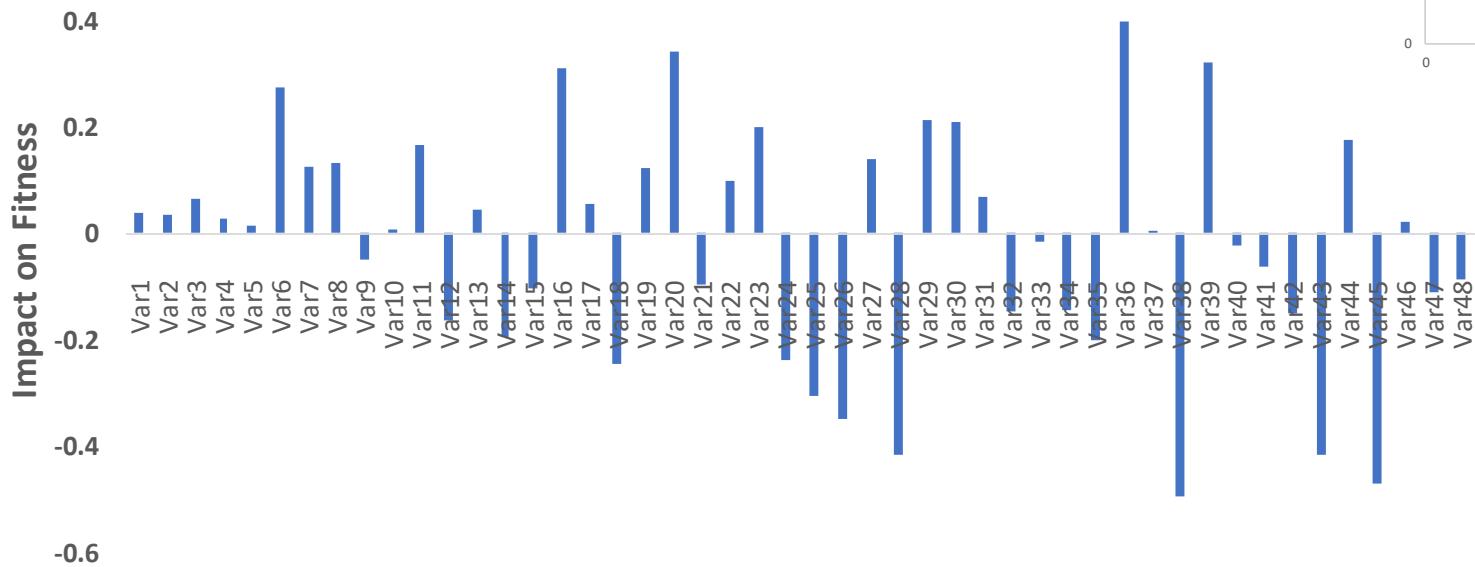
# Multi-Functional Modeling



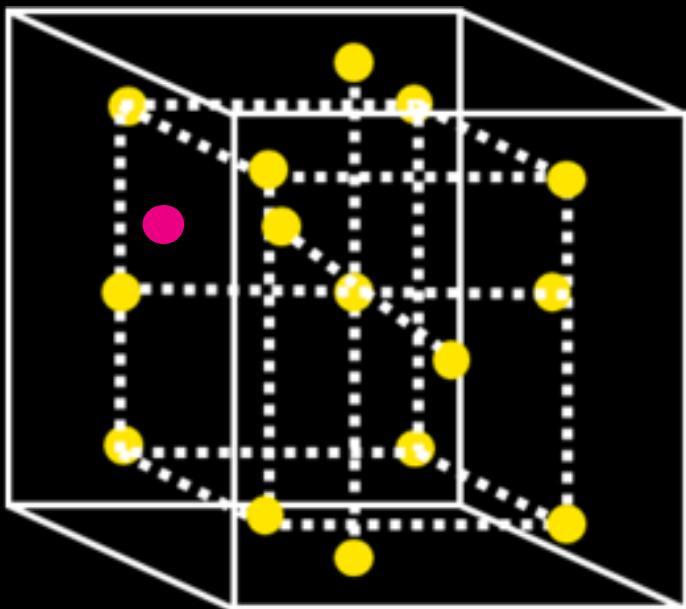
# Multi-Dimensional Fitness

Fitness =  
6x Function A  
1x Function B  
3x Function C  
5x Function D  
2x Function E  
2x Function F  
5x Function G  
5x Function H

Variations are independent substitutions or combinations



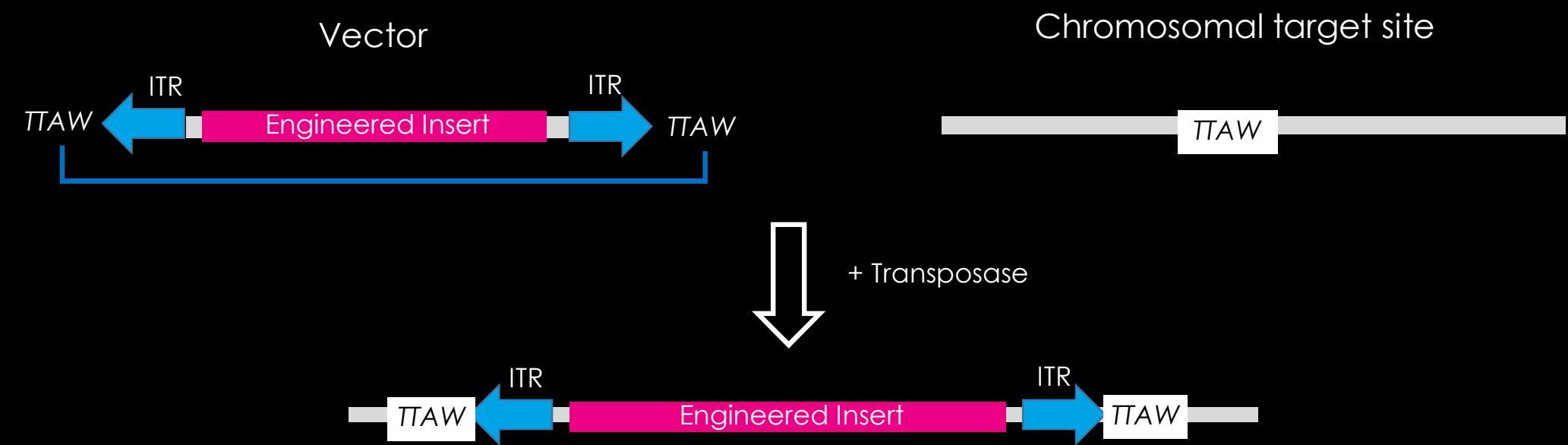
# Life is Multidimensional



## Winner

- ✓ Cell based assay
- ✓ Antigen binding
- ✓ SEC HPLC
- ✓ Tm stable
- ✓ Humanized
- ✓ 'AdiMab 7'
- ✓ ...And more

# Leap-In® Mediated Integration



- Single copy integration at each site
- Multiple insertions across the genome

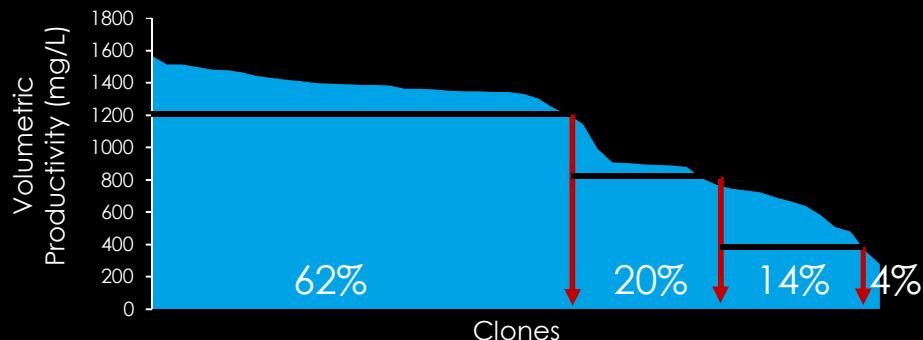
W = A or T  
2 Transposases



# Robust High Titers Fast

## High productivity stable pools

Clonal distribution in Leap-In transposase® mediated stable pools



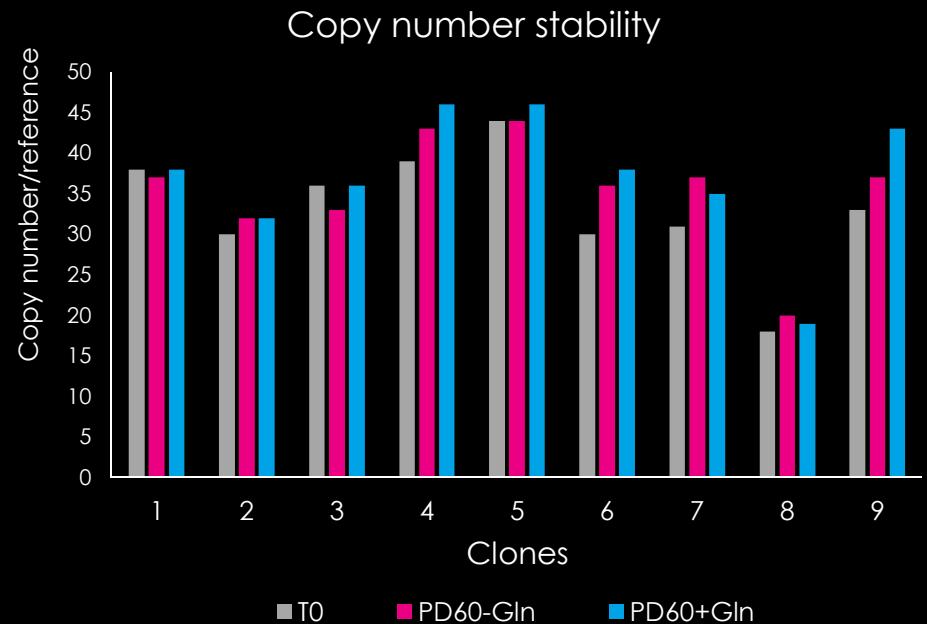
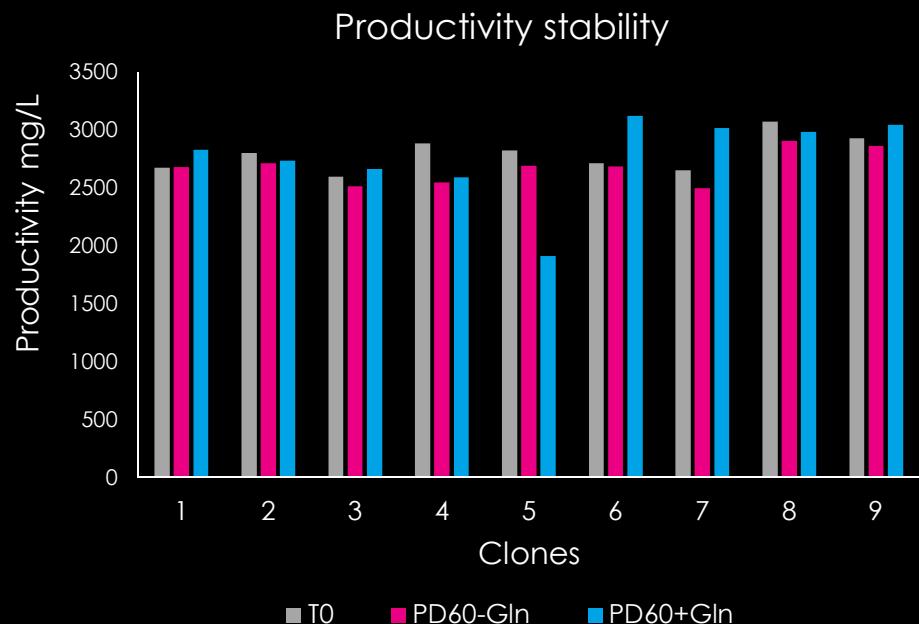
Representative Pool Titers

| Protein | Volumetric productivity | Specific productivity |
|---------|-------------------------|-----------------------|
| IgG1    | 4.2 g/L                 | 42 pcd                |
| IgG1    | 3.6 g/L                 | 29 pcd                |
| IgG1    | 3.3 g/L                 | 29 pcd                |
| IgG1    | 2.8 g/L                 | 30 pcd                |
| IgG1    | 4.2 g/L                 | 33 pcd                |
| IgG4    | 5.0 g/L                 | 43 pcd                |
| IgG4    | 5.0 g/L                 | 49 pcd                |

- 62% of clones in top quartile of expressers
- 82% of clones in top half of expressers

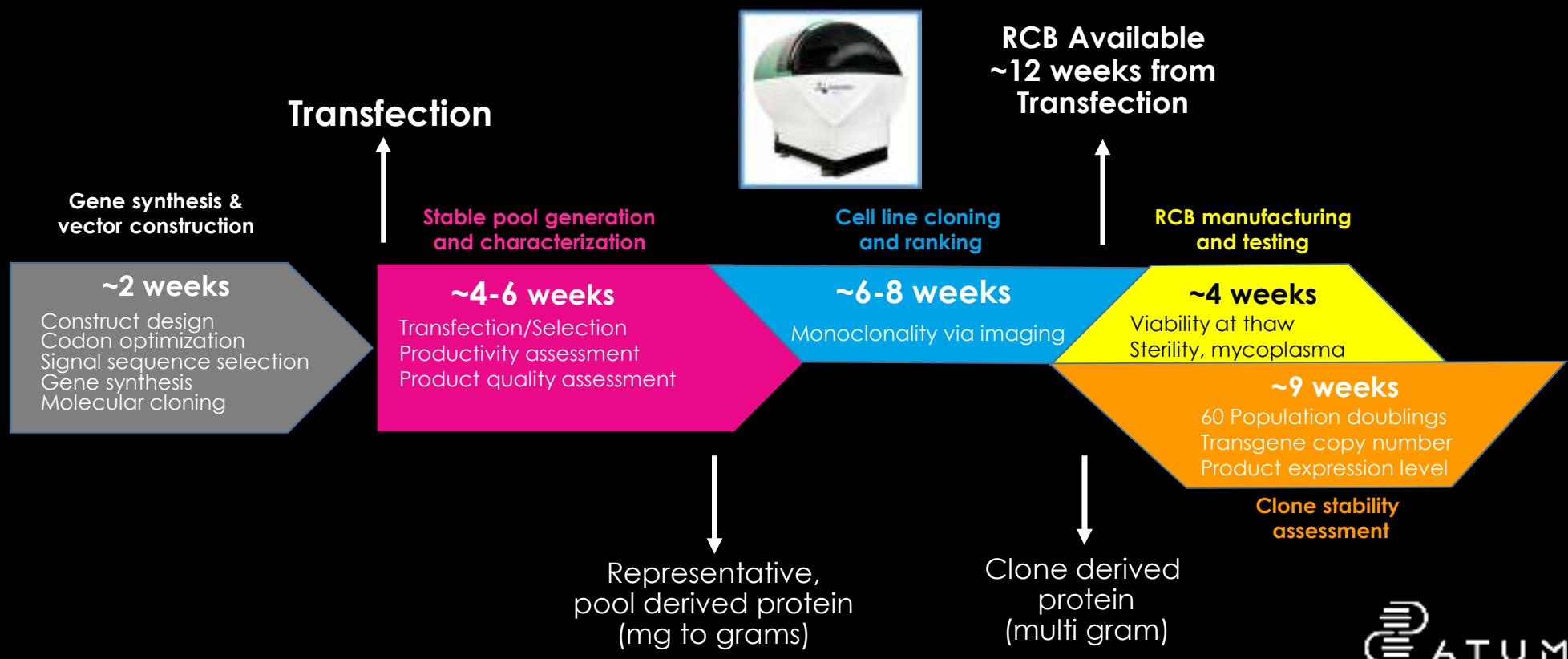


# Highly Stable Clones



# Timeline

Transfection to RCB in 12 weeks



# Thank You

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Technology presented is protected by issued US patents 10233454, 10041077, 9771402, 9580697, 9574209, 9534234, 9493521, 9428767, 9290552, 9206433, 9102944, 8975042, 8825411, 8635029, 8412461, 8401798, 8323930, 8158391, 8126653, 8005620, 7805252, 7561973, 7561972 and pending applications

